

Mon Mar 17 16:47:31 2003

US-09-840-243b-11.ra1

GenCore version 5.1.4 p5_4578
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OM protein - protein search, using sw model

Run on: March 17, 2003, 16:38:57 ; Search time 15 Seconds
(without alignments)
509.997 Million cell updates/sec

Title: US-09-840-243B-11
Perfect score: 1341
Sequence: 1 MELTOPAEDLIQTOQTASE.....VIENHILKLFQSNLVPADPE 260

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 262574 seqs, 29422922 residues
262574

Total number of hits satisfying chosen parameters:
Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Issued Patents AA: *
1: /cgn2_6/ptodata/2/iaa/5A-COMB.pep: *
2: /cgn2_6/ptodata/2/iaa/5B-COMB.pep: *
3: /cgn2_6/ptodata/2/iaa/6A-COMB.pep: *
4: /cgn2_6/ptodata/2/iaa/6B-COMB.pep: *
5: /cgn2_6/ptodata/2/iaa/6C-COMB.pep: *
6: /cgn2_6/ptodata/2/iaa/backfile1.pep: *

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Match	Length	DB	ID	Description
1	1341	100.0	260	2	US-09-172-977-1	Sequence 1, Appli
2	205	15.3	348	2	US-09-031-485-28	Sequence 28, Appli
3	205	15.3	348	2	US-08-847-429A-28	Sequence 28, Appli
4	205	15.3	348	3	US-09-065-474-28	Sequence 28, Appli
5	205	15.3	348	4	US-09-557-034-28	Sequence 33, Appli
6	205	15.3	1745	2	US-08-847-429A-33	Sequence 33, Appli
7	205	15.3	1745	3	US-09-065-474-33	Sequence 33, Appli
8	205	15.3	1745	4	US-09-557-034-33	Sequence 4, Appli
9	205	15.3	1839	2	US-09-172-977-4	Sequence 3, Appli
10	202.5	15.1	843	2	US-09-172-977-3	Sequence 2, Appli
11	197	14.7	1088	4	US-09-196-387-10	Sequence 10, Appli
12	187.5	13.9	673	4	US-09-196-387-8	Sequence 2, Appli
13	187	13.9	949	4	US-09-196-387-2	Sequence 2, Appli
14	187	13.9	1327	2	US-08-462-481-2	Sequence 2, Appli
15	187	13.9	741	2	US-08-436-771-2	Sequence 2, Appli
16	171.5	12.8	741	2	US-08-436-998-2	Sequence 2, Appli
17	171.5	12.8	741	2	US-08-487-797-2	Sequence 2, Appli
18	171.5	12.8	741	2	US-08-701-005A-2	Sequence 2, Appli
19	171.5	12.8	741	2	US-08-479-895-2	Sequence 2, Appli
20	171.5	12.8	741	2	US-08-943-956A-2	Sequence 2, Appli
21	171.5	12.8	741	5	PCT-US95-02058-2	Sequence 2, Appli
22	171.5	12.8	741	5	PCT-US95-02058-2	Sequence 2, Appli
23	171.5	12.8	741	2	US-08-436-771-4	Sequence 4, Appli
24	169.5	12.6	741	2	US-08-434-998-4	Sequence 4, Appli
25	169.5	12.6	741	2	US-08-487-797-4	Sequence 4, Appli
26	169.5	12.6	741	2	PCT-US95-02058-4	Sequence 4, Appli
27	169.5	12.6	741	5	PCT-US95-02058-4	Sequence 4, Appli

28	165.5	12.3	452	3	US-09-035-706-2	Sequence 2, Appli
29	165.5	12.3	452	3	US-08-955-841-2	Sequence 2, Appli
30	165.5	12.3	452	4	US-09-390-425-2	Sequence 2, Appli
31	165.5	12.3	452	4	US-09-566-906-2	Sequence 3, Appli
32	163	12.2	118	4	US-08-934-131-3	Sequence 2, Appli
33	161	12.0	118	3	US-08-965-904B-2	Sequence 1, Appli
34	161	12.0	118	4	US-08-934-131-1	Sequence 154, App
35	160.5	12.0	267	4	US-09-071-035-154	Sequence 156, App
36	159	11.9	238	4	US-09-071-035-156	Sequence 379, App
37	158.5	11.8	656	4	US-09-605-785-379	Sequence 379, App
38	158.5	11.8	656	4	US-09-439-313-379	Sequence 380, App
39	158.5	11.8	671	4	US-09-352-616A-379	Sequence 380, App
40	158.5	11.8	671	4	US-09-439-313-380	Sequence 380, App
41	158.5	11.8	671	4	US-09-352-616A-380	Sequence 378, App
42	158.5	11.8	671	4	US-09-605-785-378	Sequence 378, App
43	158.5	11.8	1719	4	US-09-439-313-378	Sequence 378, App
44	158.5	11.8	1719	4	US-09-439-313-378	Sequence 378, App
45	158.5	11.8	1719	4	US-09-352-616A-378	Sequence 378, App

ALIGNMENTS

RESULT 1
US-09-172-977-1
Sequence 1, Application US/09172977
Patent No. 5989863
GENERAL INFORMATION:
APPLICANT: Tang, Y. Tom
APPLICANT: Guegler, Karl J.
APPLICANT: Corley, Neil C.
APPLICANT: Yue, Henry
TITLE OF INVENTION: HUMAN ANKYRIN FAMILY PROTEIN
FILE REFERENCE: PF-0615 US
CURRENT APPLICATION NUMBER: US/09/172,977
CURRENT FILING DATE: 1998-10-14
NUMBER OF SEQ ID NOS: 4
SOFTWARE: PERL Program
SEQ ID NO 1
LENGTH: 260
TYPE: PRT
ORGANISM: Homo sapiens
FEATURE: -
OTHER INFORMATION: 1808075
US-09-172-977-1

Query Match 100.0%; Score 1341; DB 2; Length 260;
Best Local Similarity 100.0%; Pred. No. 8.5e-130;
Matches 260; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY	1	MELTOPAEDLIQTOQTASELGDPEDEAAGSDTVLSLFPCTPEPVNPEDASVSS	60
DB	1	MELTOPAEDLIQTOQTASELGDPEDEAAGSDTVLSLFPCTPEPVNPEDASVSS	60
QY	61	POAGSSIKHSTLTNRQGNVSAALPATLDSLSIHQLAQGEIDQLKEHLRKGNLVNKP	120
DB	61	POAGSSIKHSTLTNRQGNVSAALPATLDSLSIHQLAQGEIDQLKEHLRKGNLVNKP	120
QY	121	DERGFTPLIWASAFGEIETVRFLEWGAADPHILAKERESALSLASTGYTDIVGLLERD	180
DB	121	DERGFTPLIWASAFGEIETVRFLEWGAADPHILAKERESALSLASTGYTDIVGLLERD	180
QY	181	VDINIYDWNGGTPLLYAVRGNHVKCEVALLARGADLTTEADSGYTPMDLAVAGYRKVOQ	240
DB	181	VDINIYDWNGGTPLLYAVRGNHVKCEVALLARGADLTTEADSGYTPMDLAVAGYRKVOQ	240
QY	241	VIENHILKLFQSNLVPADPE 260	
DB	241	VIENHILKLFQSNLVPADPE 260	

RESULT 2
US-09-031-485-28

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; Sequence 28, Application US/09031485
; Patent No. 5824306
; GENERAL INFORMATION:
; APPLICANT: Tang, Liang
; APPLICANT: Blehm, E. Scot
; TITLE OF INVENTION: DIROFILARIA AND BRUGIA ANKYRIN
; TITLE OF INVENTION: PROTEINS, NUCLEIC ACID MOLECULES, AND
; NUMBER OF SEQUENCES: 85
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Carol Talkington Verser, Ph.D.
; ADDRESS: Heska Corporation
; STREET: 1825 Sharp Point Drive
; CITY: Fort Collins
; STATE: Colorado
; COUNTRY: USA
; ZIP: 80525
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: Windows 95
; SOFTWARE: Wordperfect for Windows, Version 7.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/031,485
; FILING DATE:
; CLASSIFICATION: 530
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US/08/847,429
; FILING DATE: 24-APR-1997
; ATTORNEY/AGENT INFORMATION:
; NAME: Verser, Carol Talkington
; REGISTRATION NUMBER: 37,459
; REFERENCE/DOCKET NUMBER: HW-5
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 970/493-7272
; TELEFAX: 970/484-9505
; INFORMATION FOR SEQ ID NO: 28:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 348 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-09-031-485-28

Query Match
Best Local Similarity 15.3%; Score 205; DB 2; Length 348;
Matches 58; Conservative 28; Mismatches 77; Indels 8; Gaps 3;

QY 75 NRQGNESALPATLDSLSIHQAQGLDQKEHLRKGDNLVKNKPDERGFTPLIWASAF 134
DB 28 NSQHSNKGES-----SASFRLAARAGNLDRLVLELRSGTD-INTCNANGLNALHLASKE 80
QY 135 GEIEIVRFLLEWAGADPHILAKERESALSLASTGTYTDIVGLLEKRDVDINITYDMNGTPL 194
DB 81 GHHEVVRRLKRAVDATRKGNLTALHSLAGQELITVTVLENGANVNVQSLNGFTPL 140
QY 195 LYAVRGNHVKCYEALLARGADLTTEADSGYTPMDLAVALGY-RKVOQVIEN 244
DB 141 YMAAQENHESVRYLLAHNANQALSTEDGFTPLAVALQGHDRVAVVLEEN 191

RESULT 3
US-08-847-429A-28
; Sequence 28, Application US/08847429A
; Patent No. 5827692
; GENERAL INFORMATION:
; APPLICANT: Tang, Liang
; APPLICANT: Blehm, E. Scot
; TITLE OF INVENTION: DIROFILARIA AND BRUGIA ANKYRIN
; TITLE OF INVENTION: PROTEINS, NUCLEIC ACID MOLECULES, AND
; NUMBER OF SEQUENCES: 85
; CORRESPONDENCE ADDRESS:

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; ADDRESSEE: Carol Talkington Verser, Ph.D.
; ADDRESS: Heska Corporation
; STREET: 1825 Sharp Point Drive
; CITY: Fort Collins
; STATE: Colorado
; COUNTRY: USA
; ZIP: 80525
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: Windows 95
; SOFTWARE: Wordperfect for Windows, Version 7.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: 435
; FILING DATE: 24-APR-1997
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Verser, Carol Talkington
; REGISTRATION NUMBER: 37,459
; REFERENCE/DOCKET NUMBER: HW-5
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 970/493-7272
; TELEFAX: 970/484-9505
; INFORMATION FOR SEQ ID NO: 28:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 348 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-08-847-429A-28

Query Match
Best Local Similarity 15.3%; Score 205; DB 2; Length 348;
Matches 58; Conservative 28; Mismatches 77; Indels 8; Gaps 3;

QY 75 NRQGNESALPATLDSLSIHQAQGLDQKEHLRKGDNLVKNKPDERGFTPLIWASAF 134
DB 28 NSQHSNKGES-----SASFRLAARAGNLDRLVLELRSGTD-INTCNANGLNALHLASKE 80
QY 135 GEIEIVRFLLEWAGADPHILAKERESALSLASTGTYTDIVGLLEKRDVDINITYDMNGTPL 194
DB 81 GHHEVVRRLKRAVDATRKGNLTALHSLAGQELITVTVLENGANVNVQSLNGFTPL 140
QY 195 LYAVRGNHVKCYEALLARGADLTTEADSGYTPMDLAVALGY-RKVOQVIEN 244
DB 141 YMAAQENHESVRYLLAHNANQALSTEDGFTPLAVALQGHDRVAVVLEEN 191

RESULT 4
US-09-065-474-28
; Sequence 28, Application US/09065474
; Patent No. 6063599
; GENERAL INFORMATION:
; APPLICANT: Tang, Liang
; APPLICANT: Blehm, E. Scot
; TITLE OF INVENTION: DIROFILARIA AND BRUGIA ANKYRIN
; TITLE OF INVENTION: PROTEINS, NUCLEIC ACID MOLECULES, AND
; NUMBER OF SEQUENCES: 171
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Carol Talkington Verser, Ph.D.
; ADDRESS: Heska Corporation
; STREET: 1825 Sharp Point Drive
; CITY: Fort Collins
; STATE: Colorado
; COUNTRY: USA
; ZIP: 80525
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: Windows 95
; SOFTWARE: Wordperfect for Windows, Version 7.0
; CURRENT APPLICATION DATA:

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APPLICATION NUMBER: US/09/065,474
FILING DATE: 24-APR-1998
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Verser, Carol Talkington
REGISTRATION NUMBER: 37,459
REFERENCE/DOCKET NUMBER: HW-5-C1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 970/493-7272
TELEFAX: 970/484-9505
INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
LENGTH: 348 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-09-065-474-28

Query Match 15.3%; Score 205; DB 3; Length 348;
Best Local Similarity 33.9%; Pred. No. 7.6e-13;
Matches 58; Conservative 28; Mismatches 77; Indels 8; Gaps 3;

QY 75 NRQGNVSALPATLDSLSIHQLAQGLDQLKEHLRKGDNLVKNKPDGFTPLIWASAF 134
DB 28 NSQHSNKGES-----SASFLRARAGNLDRLVLELLRSSTD-INTCNANGNALHLASKE 80

QY 135 GEIETVRFLENGADPHILAKERESALSLASTGTYTDIVGLLELRLVDINITYDWNNGTPL 194
DB 81 GHHEVRELKRAVDATRKGNLTALHSLAQGLIVTLVENGANVNVQSLNGFTPL 140

QY 195 LYAVRGNHVKCEALLARGADLTTEADSGYTPMDLAVAGY-RKVQOVLEN 244
DB 141 YMAAQENHESVRYLLAHNANQALSTEDGFTPLAVALQCGHDRVAVVLEEN 191

RESULT 5

US-09-557-034-28
Sequence 28, Application US/09557034
Patent No. 6365569

GENERAL INFORMATION:

APPLICANT: Tang, Liang
Blehm, E. Scot
TITLE OF INVENTION: DIROFILARIA AND BRUGIA ANKYRIN
PROTEINS, NUCLEIC ACID MOLECULES, AND
USES THEREOF

NUMBER OF SEQUENCES: 171

CORRESPONDENCE ADDRESS:

ADDRESSEE: Carol Talkington Verser, Ph.D.

STREET: 1825 Sharp Point Drive

CITY: Fort Collins

STATE: Colorado

COUNTRY: USA

ZIP: 80525

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: Windows 95

SOFTWARE: WordPerfect for Windows, Version 7.0

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/557,034

FILING DATE: 21-APR-2000

CLASSIFICATION: <Unknown>

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 09/065,474

FILING DATE: <Unknown>

ATTORNEY/AGENT INFORMATION:

NAME: Verser, Carol Talkington

REGISTRATION NUMBER: 37,459

REFERENCE/DOCKET NUMBER: HW-5-C1

TELECOMMUNICATION INFORMATION:

TELEPHONE: 970/493-7272

TELEFAX: 970/484-9505

INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
LENGTH: 348 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 28:
US-09-557-034-28

Query Match 15.3%; Score 205; DB 4; Length 348;
Best Local Similarity 33.9%; Pred. No. 7.6e-13;
Matches 58; Conservative 28; Mismatches 77; Indels 8; Gaps 3;

QY 75 NRQGNVSALPATLDSLSIHQLAQGLDQLKEHLRKGDNLVKNKPDGFTPLIWASAF 134
DB 28 NSQHSNKGES-----SASFLRARAGNLDRLVLELLRSSTD-INTCNANGNALHLASKE 80

QY 135 GEIETVRFLENGADPHILAKERESALSLASTGTYTDIVGLLELRLVDINITYDWNNGTPL 194
DB 81 GHHEVRELKRAVDATRKGNLTALHSLAQGLIVTLVENGANVNVQSLNGFTPL 140

QY 195 LYAVRGNHVKCEALLARGADLTTEADSGYTPMDLAVAGY-RKVQOVLEN 244
DB 141 YMAAQENHESVRYLLAHNANQALSTEDGFTPLAVALQCGHDRVAVVLEEN 191

RESULT 6

US-09-031-485-33
Sequence 33, Application US/09031485
Patent No. 5824306

GENERAL INFORMATION:

APPLICANT: Tang, Liang
Blehm, E. Scot
TITLE OF INVENTION: DIROFILARIA AND BRUGIA ANKYRIN
PROTEINS, NUCLEIC ACID MOLECULES, AND
USES THEREOF

NUMBER OF SEQUENCES: 85

CORRESPONDENCE ADDRESS:

ADDRESSEE: Carol Talkington Verser, Ph.D.

STREET: 1825 Sharp Point Drive

CITY: Fort Collins

STATE: Colorado

COUNTRY: USA

ZIP: 80525

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: Windows 95

SOFTWARE: WordPerfect for Windows, Version 7.0

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/031,485

FILING DATE:

CLASSIFICATION: 530

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/08/847,429

FILING DATE: 24-APR-1997

ATTORNEY/AGENT INFORMATION:

NAME: Verser, Carol Talkington

REGISTRATION NUMBER: 37,459

REFERENCE/DOCKET NUMBER: HW-5

TELECOMMUNICATION INFORMATION:

TELEPHONE: 970/493-7272

TELEFAX: 970/484-9505

INFORMATION FOR SEQ ID NO: 33:

SEQUENCE CHARACTERISTICS:

LENGTH: 1745 amino acids

TYPE: amino acid

TOPOLOGY: linear

MOLECULE TYPE: protein

Query Match 15.3%; Score 205; DB 2; Length 1745;

NUMBER OF SEQUENCES: 171
CORRESPONDENCE ADDRESS:
ADDRESSEE: Carol Talkington Verser, Ph.D.
Heska Corporation
STREET: 1825 Sharp Point Drive
CITY: Fort Collins
STATE: Colorado
COUNTRY: USA
ZIP: 80525
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: Windows 95
SOFTWARE: Wordperfect for windows, Version 7.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/557,034
FILING DATE: 21-Apr-2000
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/065,474
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: Verser, Carol Talkington
REGISTRATION NUMBER: 37,459
REFERENCE/DOCKET NUMBER: HM-5-C1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 970/493-7272
TELEFAX: 970/484-9505
INFORMATION FOR SEQ ID NO: 33:
SEQUENCE CHARACTERISTICS:
LENGTH: 1745 amino acids
TYPE: amino acid
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 33:
US-09-557-034-33

Query Match 15.3%; Score 205; DB 4; Length 1745;
Best Local Similarity 33.9%; Pred. No. 9.4e-12;
Matches 58; Conservative 28; Mismatches 77; Indels 8; Gaps 3;

QY 75 NRQGNVSALPATLDSLSHQLAQGLDQKHLRKGDNLVKNKPDGFTPLIWASAF 134
DB 28 NSQHSNKGES-----SASFRAARAGNLDRLVLELRSGTD-INTCNANGINALHLASKE 80
QY 135 GEIETVRFLEMGADPHILAKRESALSLASTGGYTDIVGLLERVDVINIYDWMNGTPL 194
DB 81 GHHEVRELKRRKADVDATRKGNLTALHIASLAGQELIVTVLVENGANVNVQSLNGFTPL 140
QY 195 LYAVRGNHVKVEALLARGADLTTEADSGYTPMDLAVAGY-RKYQOVIEEN 244
DB 141 YMAAQENHESVVRYLHMANQALSTEDGFTPLAVALQGGHDRVAVLLEN 191

RESULT 10
US-09-172-977-4
Sequence 4, Application US/09172977
Patent No. 5989863
GENERAL INFORMATION:
APPLICANT: Tang, Y. Tom
APPLICANT: Guegler, Karl J.
APPLICANT: Corley, Neil C.
TITLE OF INVENTION: HUMAN ANKYRIN FAMILY PROTEIN
FILE REFERENCE: PF-0615 US
CURRENT APPLICATION NUMBER: US/09/172,977
NUMBER OF SEQ ID NOS: 4
SOFTWARE: PERL Program
SEQ ID NO 4
LENGTH: 1839
TYPE: PRT
ORGANISM: Homo sapiens

FEATURE:
OTHER INFORMATION: 929491
US-09-172-977-4

Query Match 15.1%; Score 202.5; DB 2; Length 1839;
Best Local Similarity 29.3%; Pred. No. 1.8e-11;
Matches 56; Conservative 43; Mismatches 81; Indels 11; Gaps 4;

QY 55 DASVSSPQAGSLKSTTLTNRQGNVSALPATLDSLSHQLAQGLDQKHLRKGD 114
DB 5 DAAQKS-DSGEKFNSSQRRKRPKSDSNA-----SFLRAARAGNLDKVEYLVKGI 55
QY 115 NLVKNKPDGFTPLIWASAFGEIETVRFLEMGADPHILAKRESALSLASTGGYTDIVG 174
DB 56 D-INTCNQNGINALHLAAKEGHVGLVQELLGRGSSVDSATKGNLTALHIASLAGQAEVVK 114
QY 175 LLERVDVINIYDWMNGTPLLVAVRGNHVKVEALLARGADLTTEADSGYTPMDLAVAG 234
DB 115 VLVKEGANINAQSQNGFTPLYMAAQENHIDVVKYLLENGANQSTATEDGFTPLAVALQGG 174
QY 235 YRK-VQOVIEEN 244
DB 175 HNOQVAIILEN 185

RESULT 11
US-09-172-977-3
Sequence 3, Application US/09172977
Patent No. 5989863
GENERAL INFORMATION:
APPLICANT: Tang, Y. Tom
APPLICANT: Guegler, Karl J.
APPLICANT: Corley, Neil C.
TITLE OF INVENTION: HUMAN ANKYRIN FAMILY PROTEIN
FILE REFERENCE: PF-0615 US
CURRENT APPLICATION NUMBER: US/09/172,977
NUMBER OF SEQ ID NOS: 4
SOFTWARE: PERL Program
SEQ ID NO 3
LENGTH: 843
TYPE: PRT
ORGANISM: Rattus norvegicus
FEATURE:
OTHER INFORMATION: g1841966
US-09-172-977-3

Query Match 14.7%; Score 197; DB 2; Length 843;
Best Local Similarity 32.4%; Pred. No. 2e-11;
Matches 48; Conservative 35; Mismatches 63; Indels 2; Gaps 2;

QY 98 AAQGLDQKHLRKGDNLVKNKPDGFTPLIWASAFGEIETVRFLEMGADPHILAKER 157
DB 4 ARAGNLDKVEYLVKGIID-INTCNQNGINALHLAAKEGHVGLVQELLGRGSSVDSATKKG 62
QY 158 ESALSLASTGGYTDIVGLLERVDVINIYDWMNGTPLLVAVRGNHVKVEALLARGADLT 217
DB 63 NTLHIASLAGQAEVVKVLVKEGANINAQSQNGFTPLYMAAQENHIDVVKYLLENGANQS 122
QY 218 TEADSGYTPMDLAVAGYRK-VQOVIEEN 244
DB 123 TATEDGFTPLAVALQGGHNOQVAIILEN 150

RESULT 12
US-09-082-059-2
Sequence 2, Application US/09082059A
Patent No. 6225086
GENERAL INFORMATION:
APPLICANT: Morrow, Jon S.
APPLICANT: Devarajan, Prasad
TITLE OF INVENTION: No. 6225086e1 Ankyrin Proteins and a Method for Their Identificat

FILE REFERENCE: 44574-5002-US
CURRENT APPLICATION NUMBER: US/09/082,059A
CURRENT FILING DATE: 1998-05-21
EARLIER APPLICATION NUMBER: 60/047356
EARLIER FILING DATE: 1997-05-21
NUMBER OF SEQ ID NOS: 19
SOFTWARE: Patent In Ver. 2.0
SEQ ID NO 2
LENGTH: 1088
TYPE: PRT
ORGANISM: Homo sapiens
US-09-082-059-2

Query Match 14.0%; Score 187.5; DB 4; Length 1088;
Best Local Similarity 25.5%; Pred. No. 2.8e-10;
Matches 62; Conservative 44; Mismatches 76; Indels 61; Gaps 6;

QY 62 QAGSSLKST-----LTNRQGNVSALPATLD---SLST-----HQLAAGE 102
DB 141 QGASPNAAATSGYTPHLHSAAREGHEDVAAF--LLDHGASLSITTKGFTPLHVAAGYK 198
QY 103 LDQLKEHLRKGDNLVKNPD-----ERGFT 126
DB 199 LEVANLLLOKSAS---PDAGKSGLTPLHVAAHYDNQKVALLLDQASPHAAKNGYT 254
QY 127 PLIWASAFGEIETVRFLEWGADEPHILAKERESALSLASTGYTDIVGLLERVDINITY 186
DB 255 PLHIAAKNQMDIATTLLEYGADANAVTRQGIASVHLAAQEGHVDWVSLLRNANVLS 314
QY 187 DWNGTPLLVAVRGNHVKCVALLARGADLTTEADSGYTPMDLAVAGYRKVQOVIEHNI 246
DB 315 NKSGLTPLHLAAQEDRVNAEVLVNGAHVDAQTKMGYTPHVGCHYGNIKIVNLLQHS 374
QY 247 LKL 249
DB 375 AKV 377

RESULT 13
US-09-196-387-8
Sequence 8, Application US/09196387
Patent No. 6277613
GENERAL INFORMATION:
APPLICANT: de Lange, Titia
APPLICANT: Smith, Susan
TITLE OF INVENTION: A PROTEIN THAT BINDS TO TRF1 AND METHODS
TITLE OF INVENTION: OF USE THEREOF
NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
ADDRESSEE: Klauber & Jackson
STREET: 411 Hackensack Avenue, 4th Floor
CITY: Hackensack
STATE: New Jersey
COUNTRY: USA
ZIP: 07601
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent In Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/196,387
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/095,225
FILING DATE: June 10, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Jackson Esq., David A.
REGISTRATION NUMBER: 26,742
REFERENCE/DOCKET NUMBER: 600-1-230 CIP1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-487-5800

TELEFAX: 201-343-1684
TELEX: 133521
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
LENGTH: 673 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-09-196-387-8

Query Match 13.9%; Score 187; DB 4; Length 673;
Best Local Similarity 24.3%; Pred. No. 1.5e-10;
Matches 72; Conservative 36; Mismatches 112; Indels 76; Gaps 9;

QY 18 ASELDPEDEPGEAAGSDTV-VLSLPCTPEPVNPEPDASVSSPQAGSSLKSTLTNR 76
DB 137 SSSSSPSPGSSSLAESPEAGVSTAPLGPAGAP---GTGVPAVSGALRE---LLEA 189
QY 77 QRGNEVSALPATLDSLSI-----HQLAAGELDQLKEHLRKGDNLVKNPDER 123
DB 190 CRNGDVSRVKRLVDAANVNAKDMAGRKSSPLHFAAGFRKDVVEHLLQMGAN-VHARDG 248
QY 124 GFTPLIWASAFGEIETVRFLL-----EM-----GADP 150
DB 249 GLIPLHNACSFGEAEVSVLLCQGADPNARDNNTPLHEAIIKIDVICIVLLQHADP 308
QY 151 HILAKERESALSAS-----TGGY-----TDIVGLLERVDINITYDN 189
DB 309 NIRNTDGSALDLADPSAKAVLTGEYKDELLEAARSNGNEEKMALLTPLNVNCHASDGR 368
QY 190 GGTPLLVAVRGNHVKCVALLARGADLTTEADSGYTPMDLAVAGYRKVQOVIEHNI 245
DB 369 KSTPLHLAAGYNRRIVQLLQHGADVHAKDKGLVPLHNACSYGHYEVELLKH 424

RESULT 14
US-09-196-387-10
Sequence 10, Application US/09196387
Patent No. 6277613
GENERAL INFORMATION:
APPLICANT: de Lange, Titia
APPLICANT: Smith, Susan
TITLE OF INVENTION: A PROTEIN THAT BINDS TO TRF1 AND METHODS
TITLE OF INVENTION: OF USE THEREOF
NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
ADDRESSEE: Klauber & Jackson
STREET: 411 Hackensack Avenue, 4th Floor
CITY: Hackensack
STATE: New Jersey
COUNTRY: USA
ZIP: 07601
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent In Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/196,387
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/095,225
FILING DATE: June 10, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Jackson Esq., David A.
REGISTRATION NUMBER: 26,742
REFERENCE/DOCKET NUMBER: 600-1-230 CIP1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-487-5800
TELEFAX: 201-343-1684
TELEX: 133521
INFORMATION FOR SEQ ID NO: 10:

SEQUENCE CHARACTERISTICS:
LENGTH: 949 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-09-196-387-10

Query Match 13.9%; Score 187; DB 4; Length 949;
Best Local Similarity 24.3%; Pred. No. 2.6e-10;
Matches 72; Conservative 36; Mismatches 112; Indels 76; Gaps 9;

QY 18 ASGLGDPEDPGEAAGSDTV-VLSLPCTPEFVNPEPDASVSSPQAGSSLKSTLTNR 76
DB 137 SSSSSSPSSPSSSLAESPEAGVSTAPLPGAAGP---GTGVPAVSGALRE--LLEA 189
QY 77 QRGNEVSALPATLDSLSI-----HQLAOGELDQLKEHLRKGDNLVKNKPD 123
DB 190 CRNGDVSRVKRLVDAANVNAMKMGARKSSPLHFAAGFGKRDVVEHLLQMGAN-VHARD 248
QY 124 GFTPLIWASAFGEIETVRFL-----EW-----GADP 150
DB 249 GLIPLHNACSFGEAEVSVLLCOGADPNARDNNWYTPLEHAIAIKKIDVCIVLLQHGADP 308
QY 151 HILAKERESALSLAS-----TGGY-----TDIVGLLERDNDINIDWN 189
DB 309 NIRNTDGKSLDLADPSAKAVLTGEYKDELLEAARSGNEEKLMLLTPLNVNCHASDGR 368
QY 190 GGTPLLYAVRGNHVCVEALLARGADLTTEADSGYTPMDLVALGYRKVQVIEHN 245
DB 369 KSTPLHLAAGVNRVRIQVLLQHGADVHAKDKGGLVPLHNACSYGHYEVTLELLKH 424

RESULT 15
US-09-196-387-2
Sequence 2, Application US/09196387
Patent No. 6277613

GENERAL INFORMATION:
APPLICANT: de Lange, Titia
APPLICANT: Smith, Susan

TITLE OF INVENTION: A PROTEIN THAT BINDS TO TRF1 AND METHODS
NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
ADDRESSEE: Klauber & Jackson
STREET: 411 Hackensack Avenue, 4th floor
CITY: Hackensack
STATE: New Jersey
COUNTRY: USA
ZIP: 07601

COMPUTER READABLE FORM:
MEDIUM TYPE: floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent in Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/196,387
FILING DATE:

CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/095,225
FILING DATE: June 10, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Jackson Esq., David A.
REGISTRATION NUMBER: 26,742
REFERENCE/DOCKET NUMBER: 600-1-230 CIP1
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-487-5800
TELEFAX: 201-343-1684
TELEX: 133521

INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 1327 amino acids
TYPE: amino acid

STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
HYPOTHETICAL: NO
US-09-196-387-2

Query Match 13.9%; Score 187; DB 4; Length 1327;
Best Local Similarity 24.3%; Pred. No. 4.3e-10;
Matches 72; Conservative 36; Mismatches 112; Indels 76; Gaps 9;

QY 18 ASGLGDPEDPGEAAGSDTV-VLSLPCTPEFVNPEPDASVSSPQAGSSLKSTLTNR 76
DB 137 SSSSSSPSSPSSSLAESPEAGVSTAPLPGAAGP---GTGVPAVSGALRE--LLEA 189
QY 77 QRGNEVSALPATLDSLSI-----HQLAOGELDQLKEHLRKGDNLVKNKPD 123
DB 190 CRNGDVSRVKRLVDAANVNAMKMGARKSSPLHFAAGFGKRDVVEHLLQMGAN-VHARD 248
QY 124 GFTPLIWASAFGEIETVRFL-----EW-----GADP 150
DB 249 GLIPLHNACSFGEAEVSVLLCOGADPNARDNNWYTPLEHAIAIKKIDVCIVLLQHGADP 308
QY 151 HILAKERESALSLAS-----TGGY-----TDIVGLLERDNDINIDWN 189
DB 309 NIRNTDGKSLDLADPSAKAVLTGEYKDELLEAARSGNEEKLMLLTPLNVNCHASDGR 368
QY 190 GGTPLLYAVRGNHVCVEALLARGADLTTEADSGYTPMDLVALGYRKVQVIEHN 245
DB 369 KSTPLHLAAGVNRVRIQVLLQHGADVHAKDKGGLVPLHNACSYGHYEVTLELLKH 424

Search completed: March 17, 2003, 16:41:31
Job time : 21 secs

